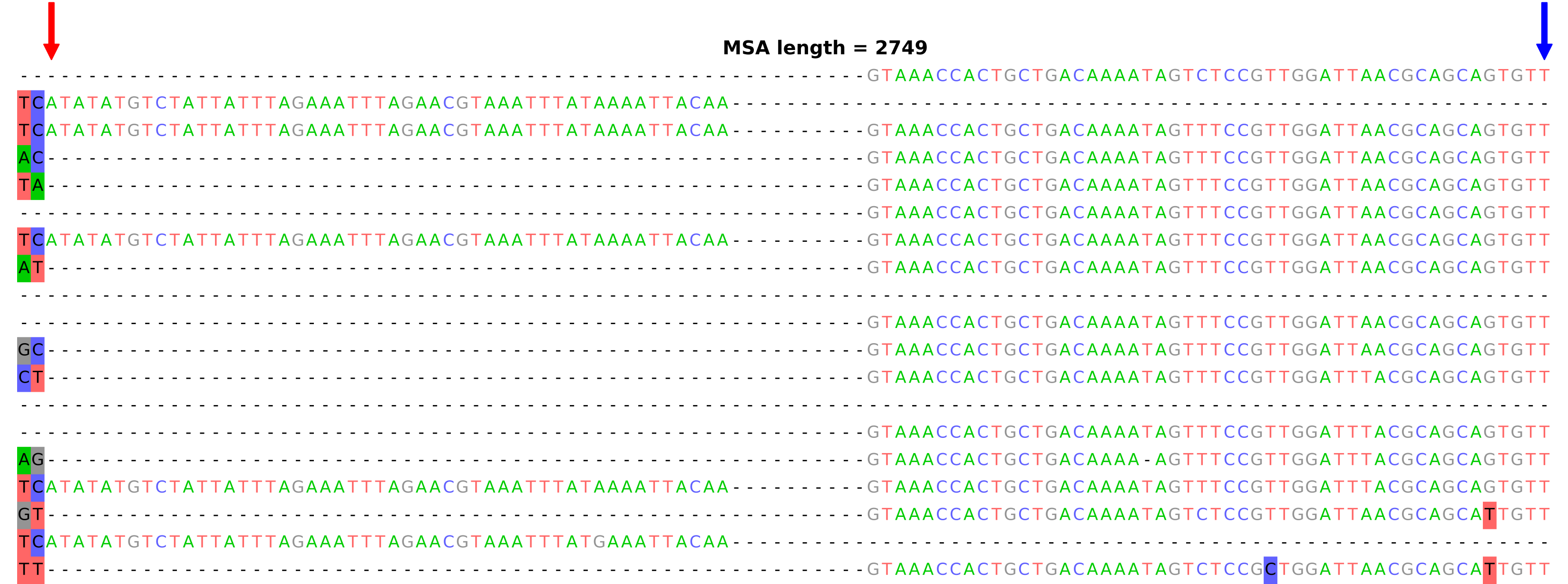
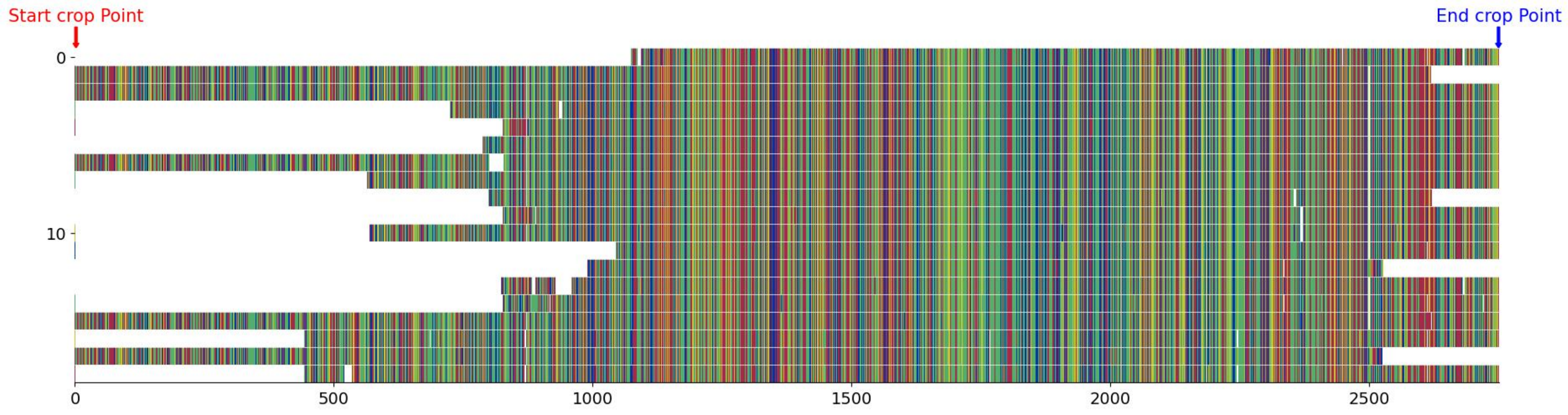


Start crop Point

End crop Point

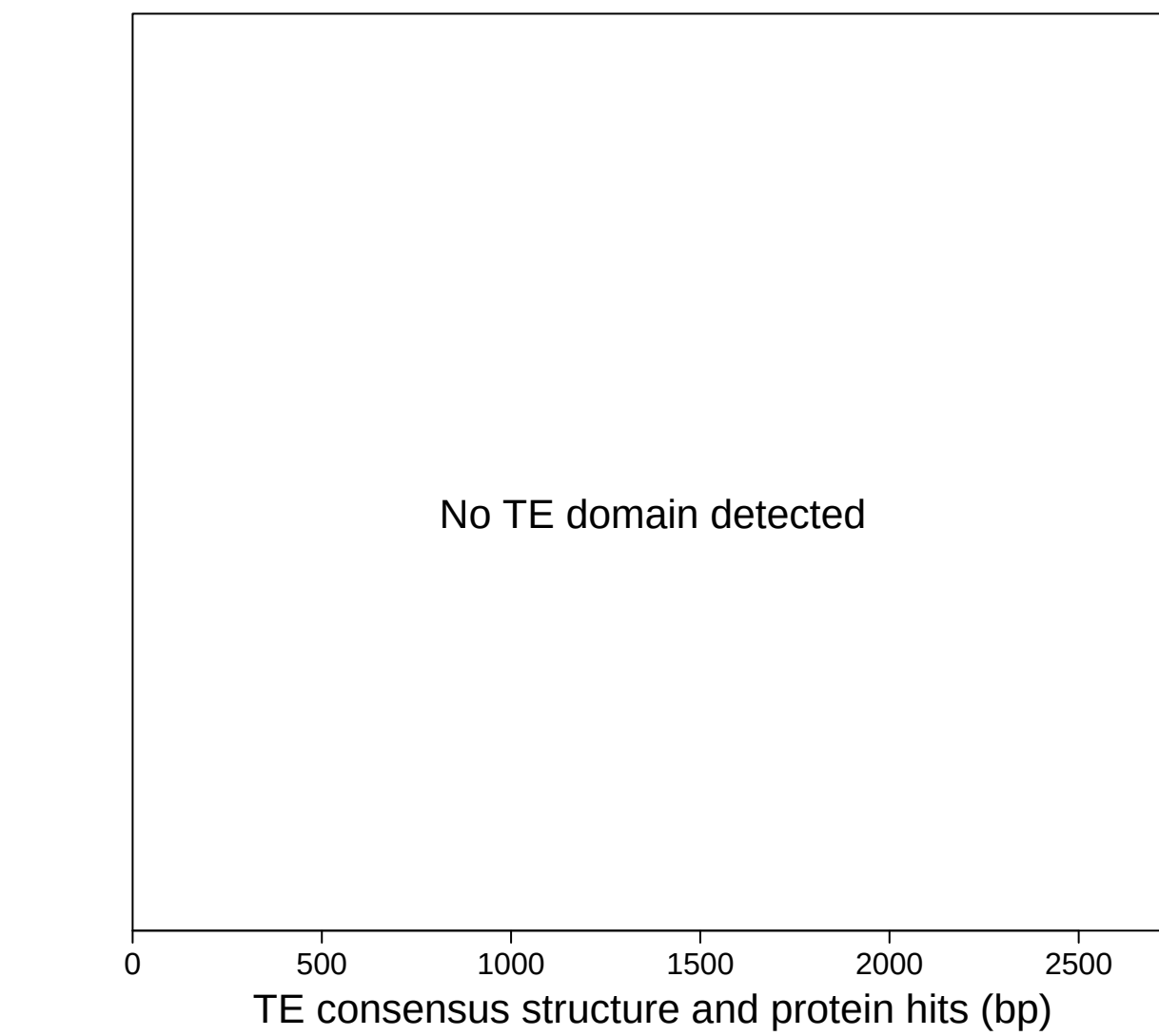
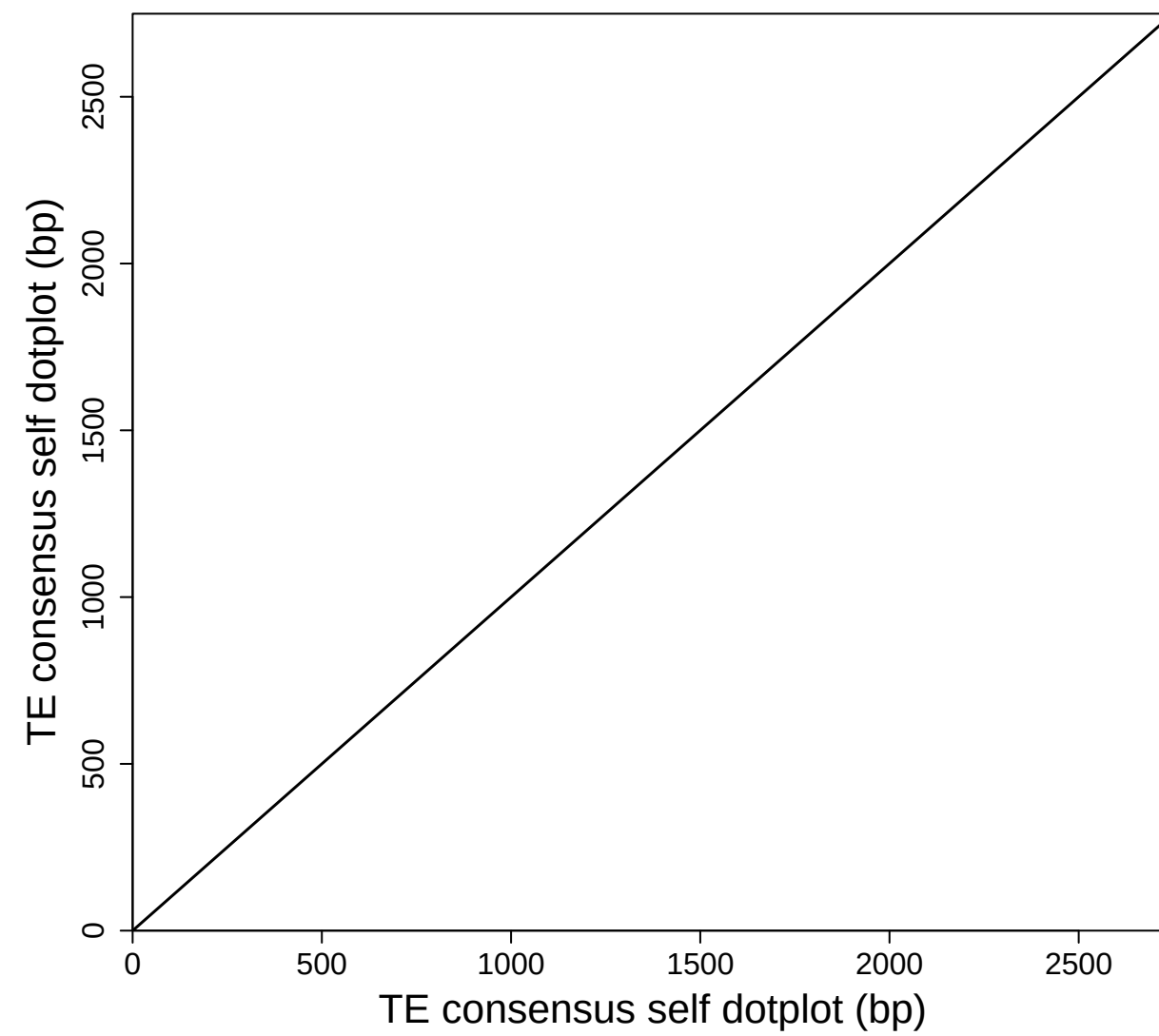
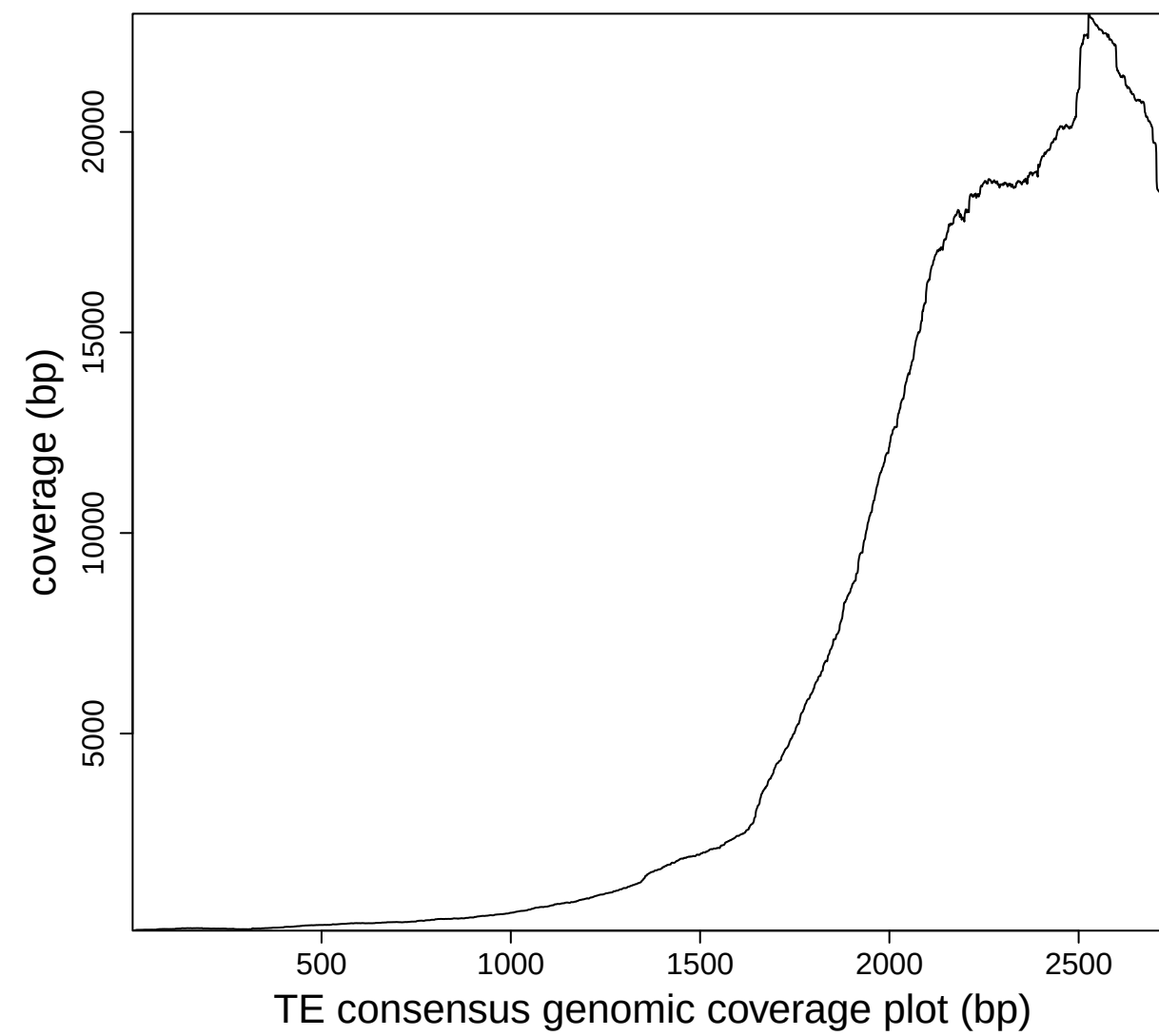
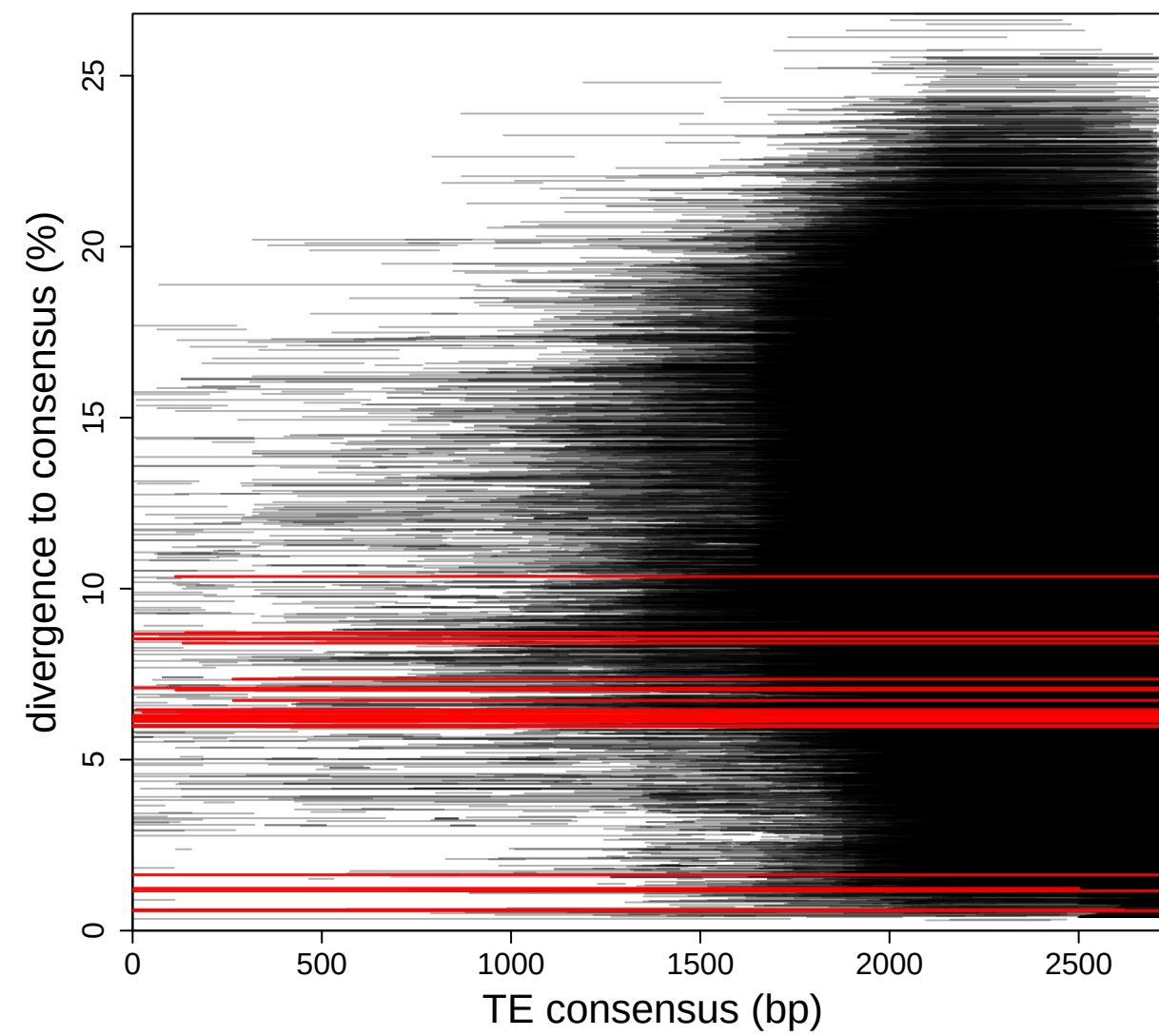
MSA length = 2749





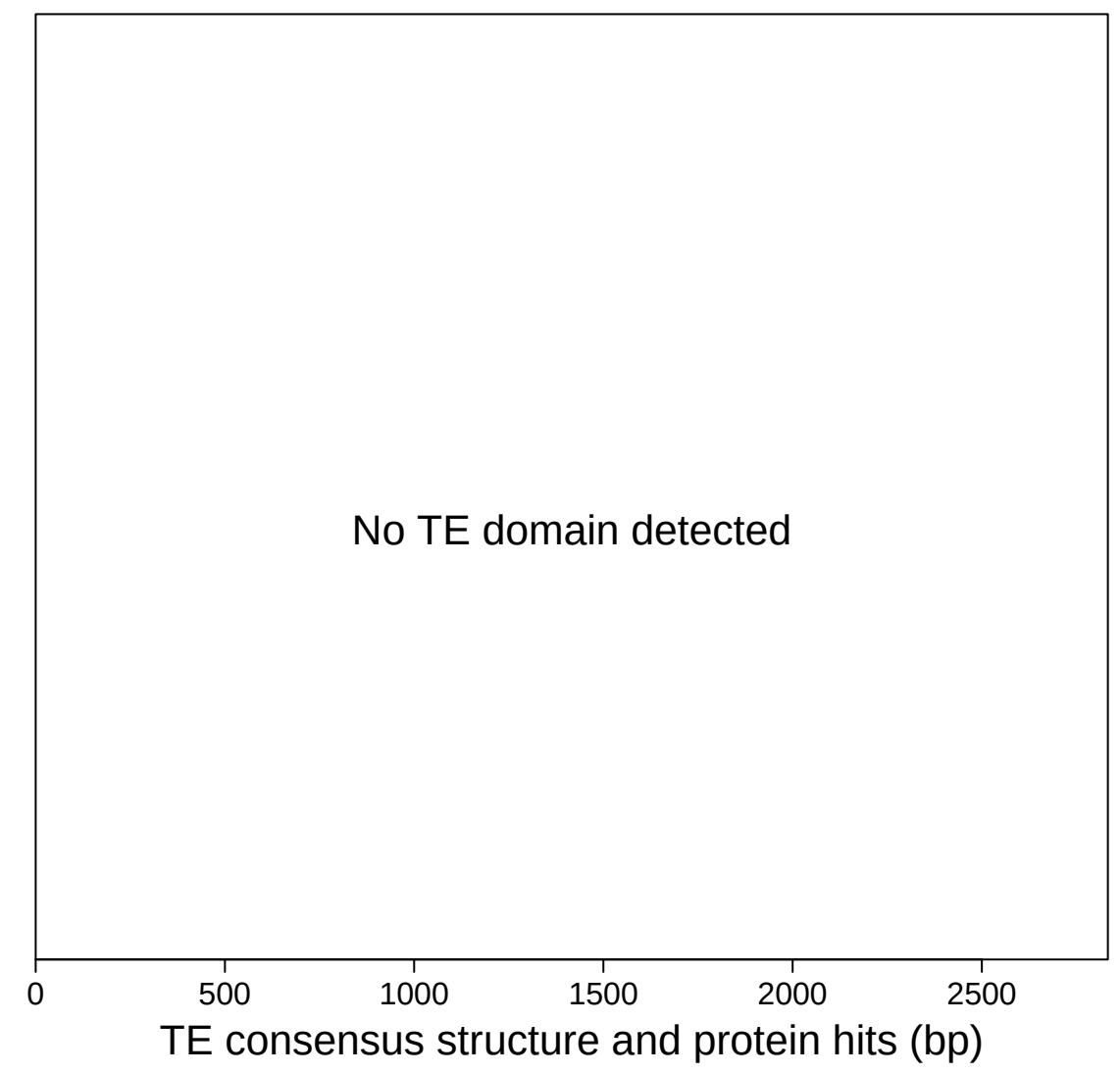
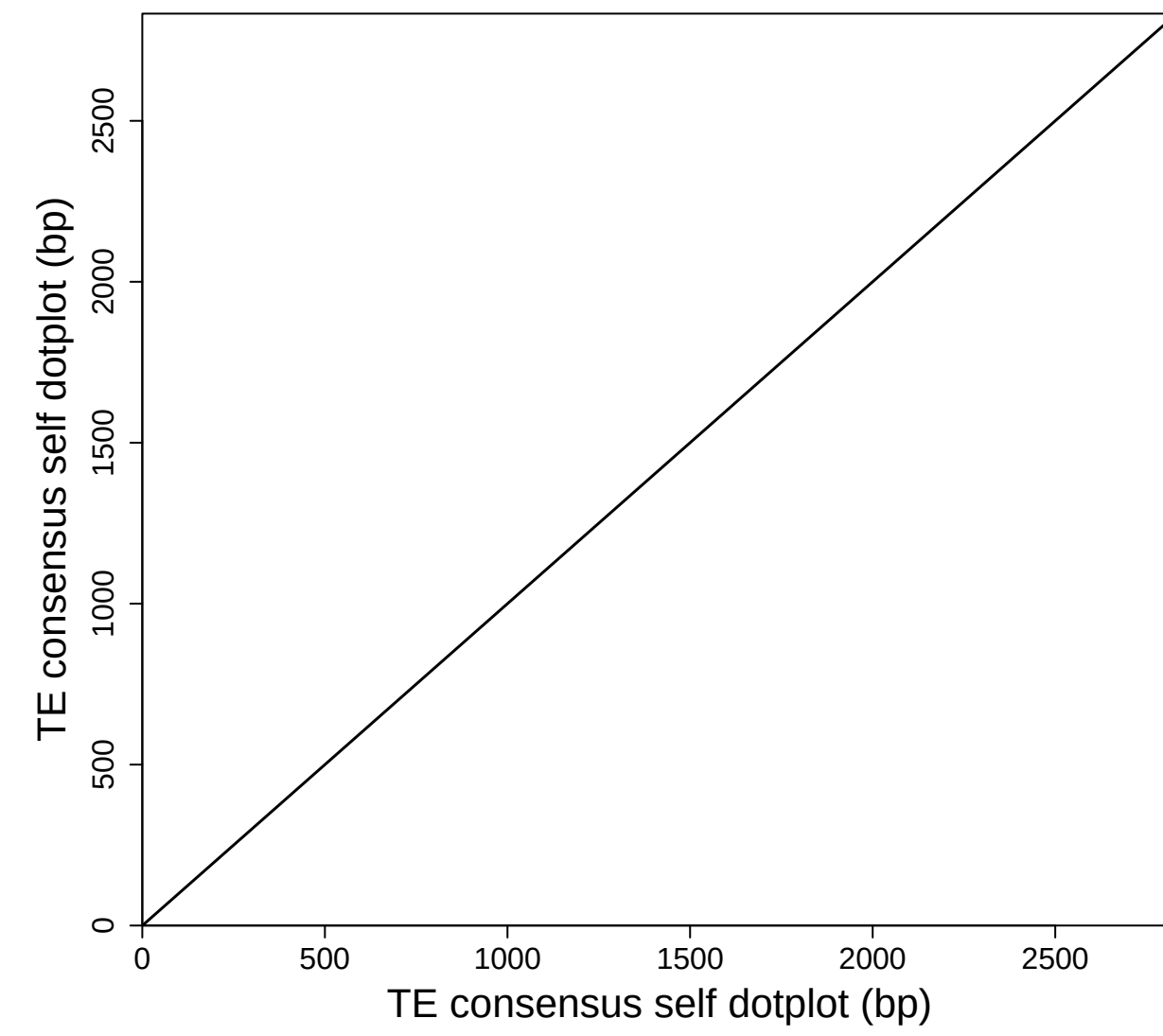
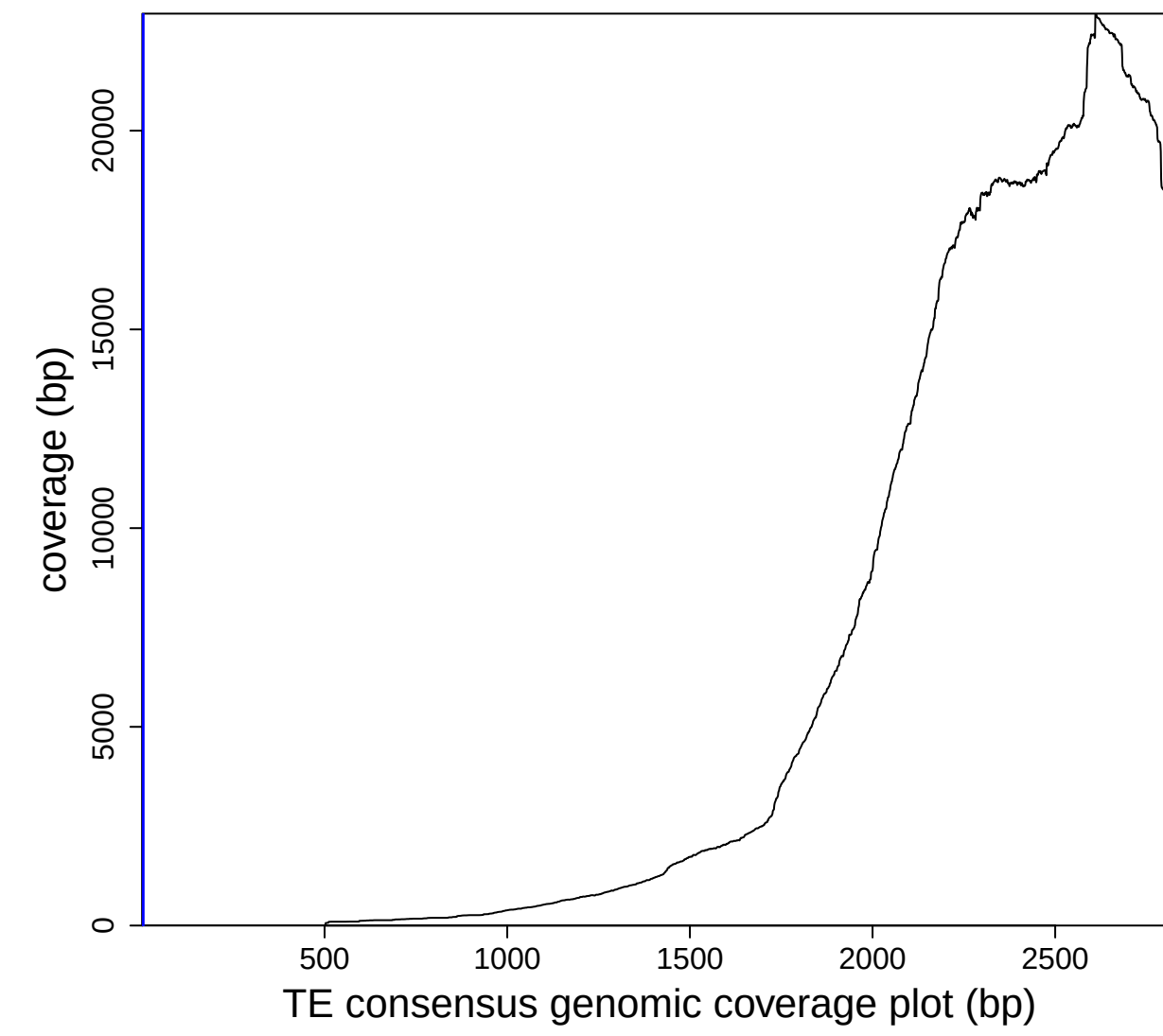
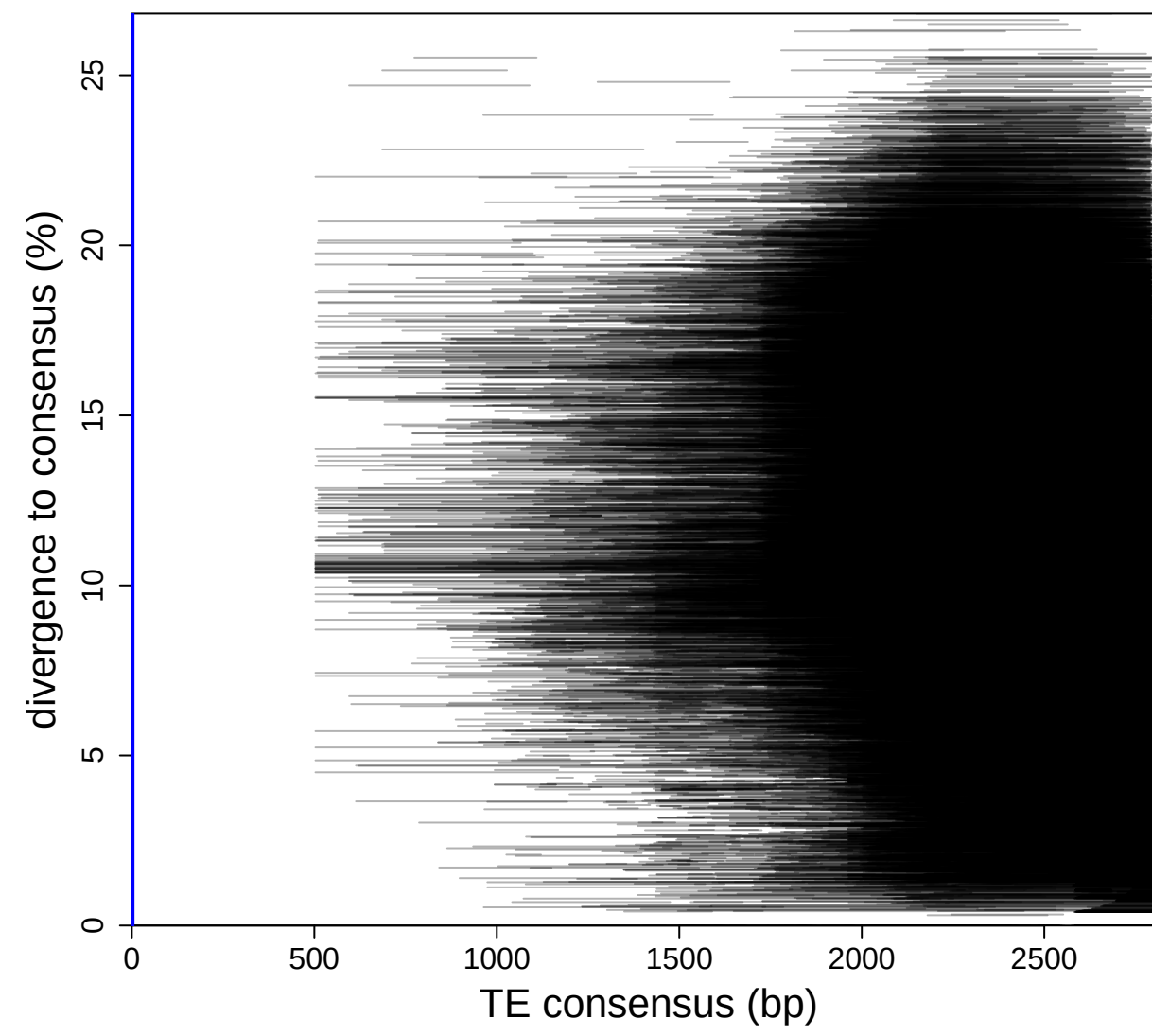
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 2749bp; fragments: 37803; full length: 33 (≥ 2474.1 bp)

After TEtrimmer 2749 bp



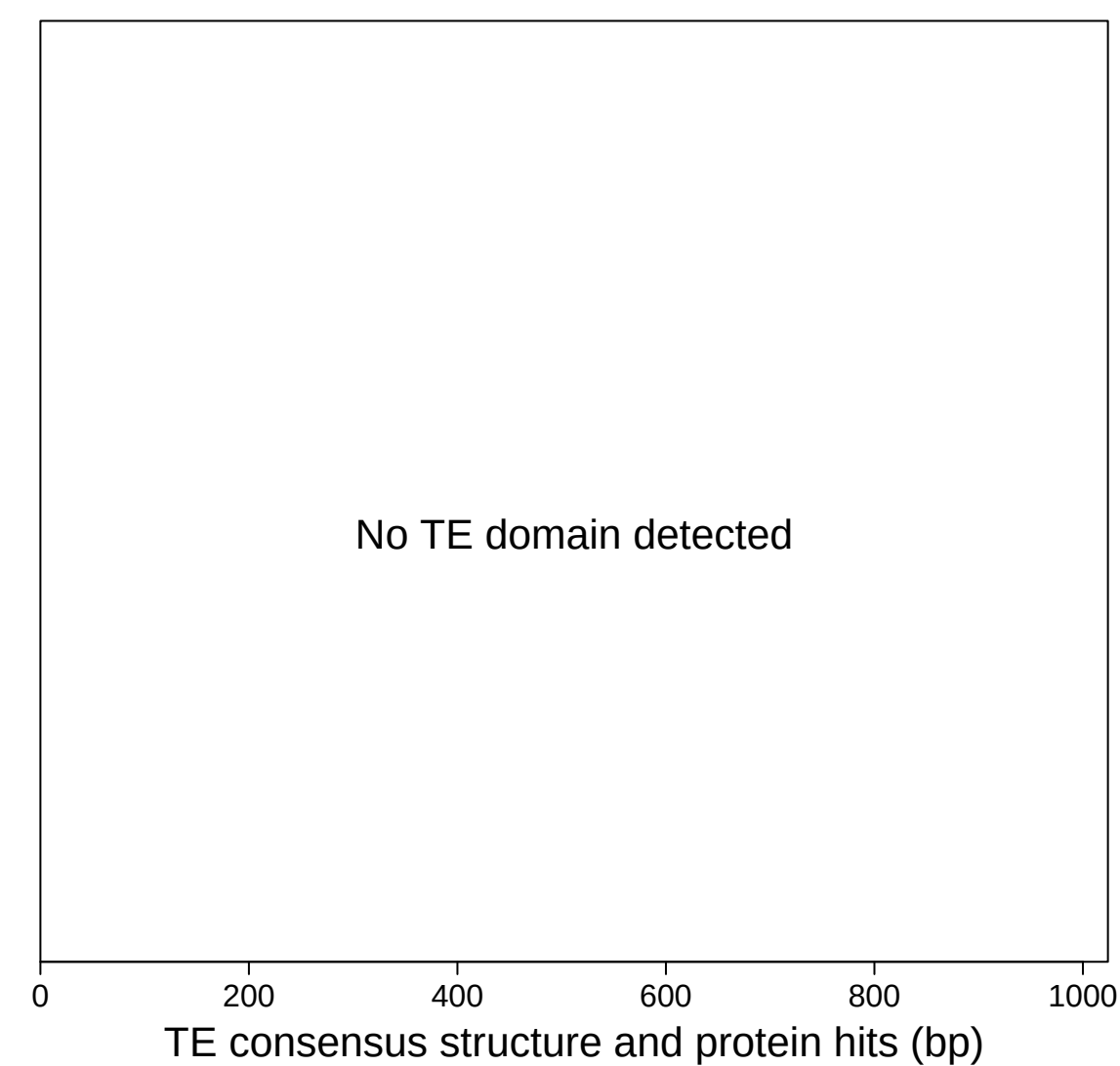
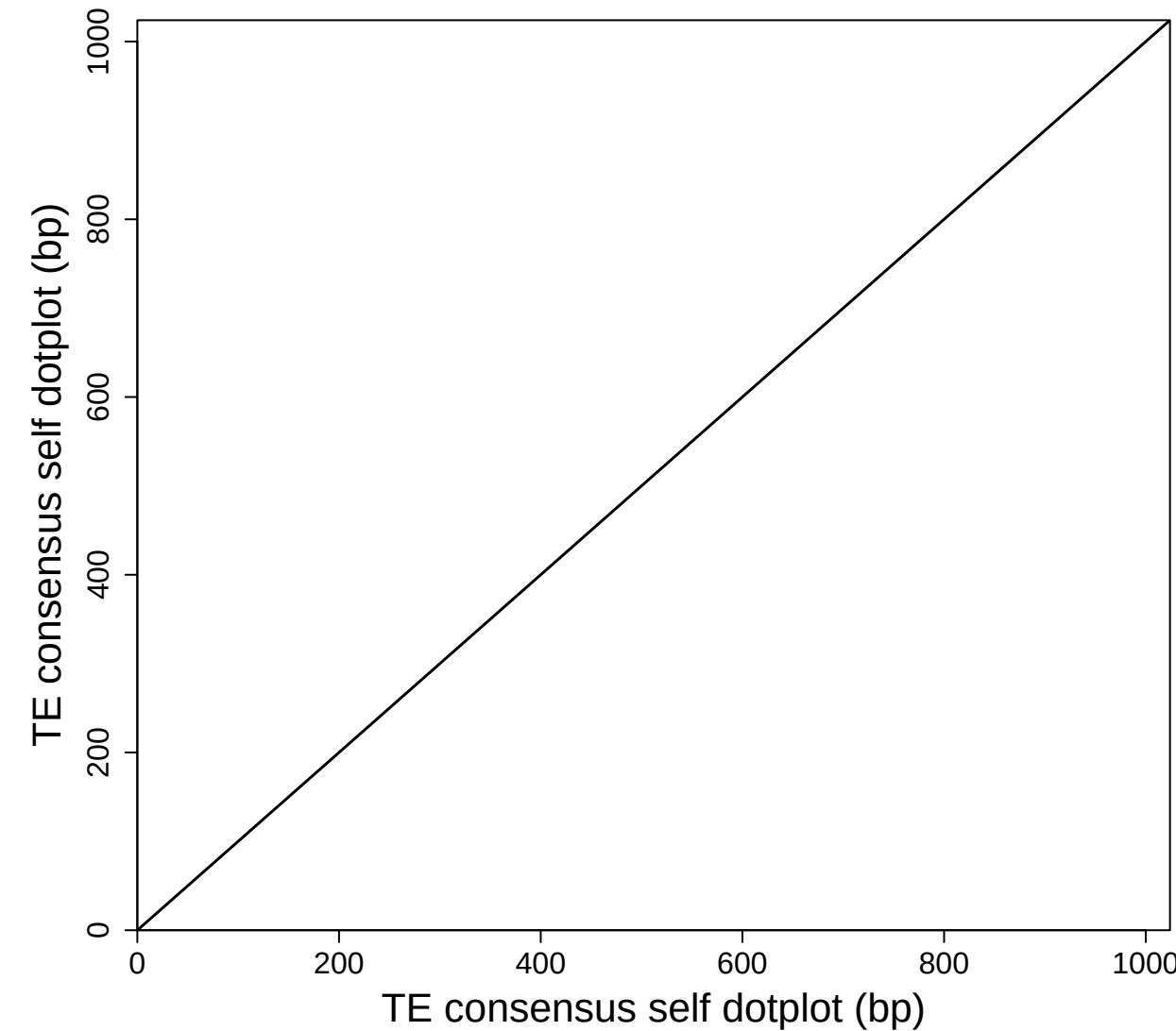
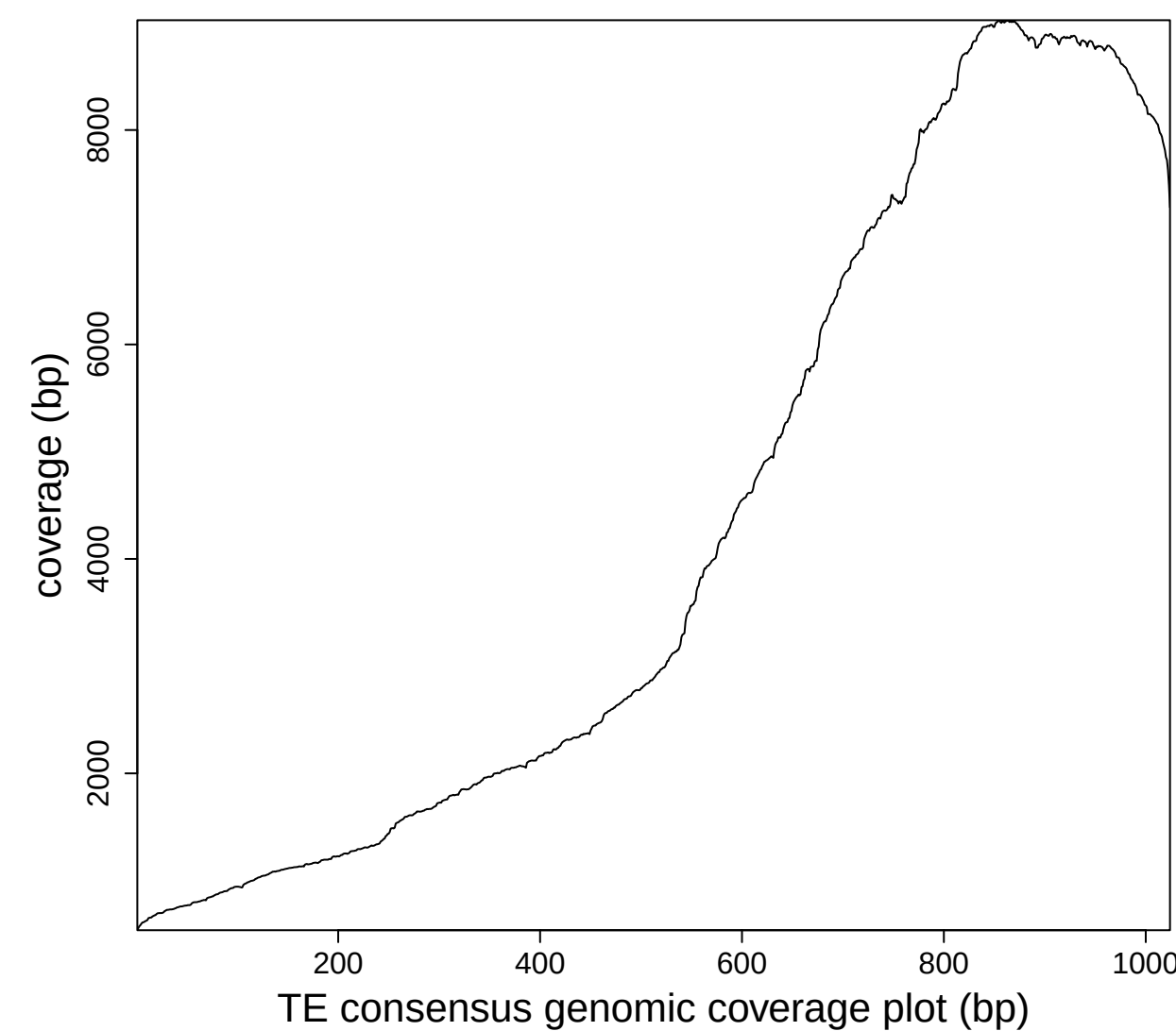
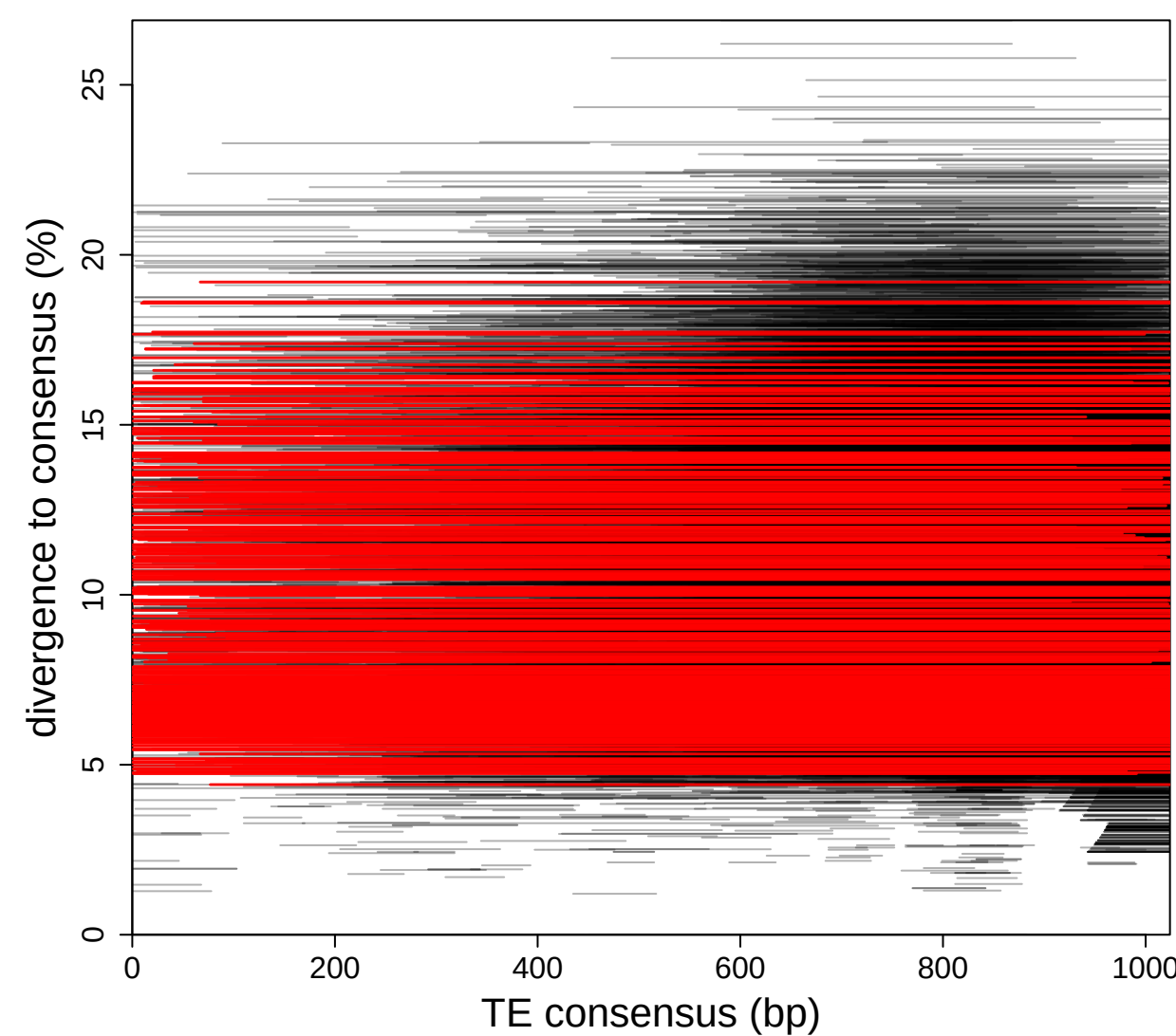
2.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 2833bp; fragments: 37441; full length: 0 (≥ 2549.7 bp)

After TEtrimmer Extended plot Blue lines are boundaries

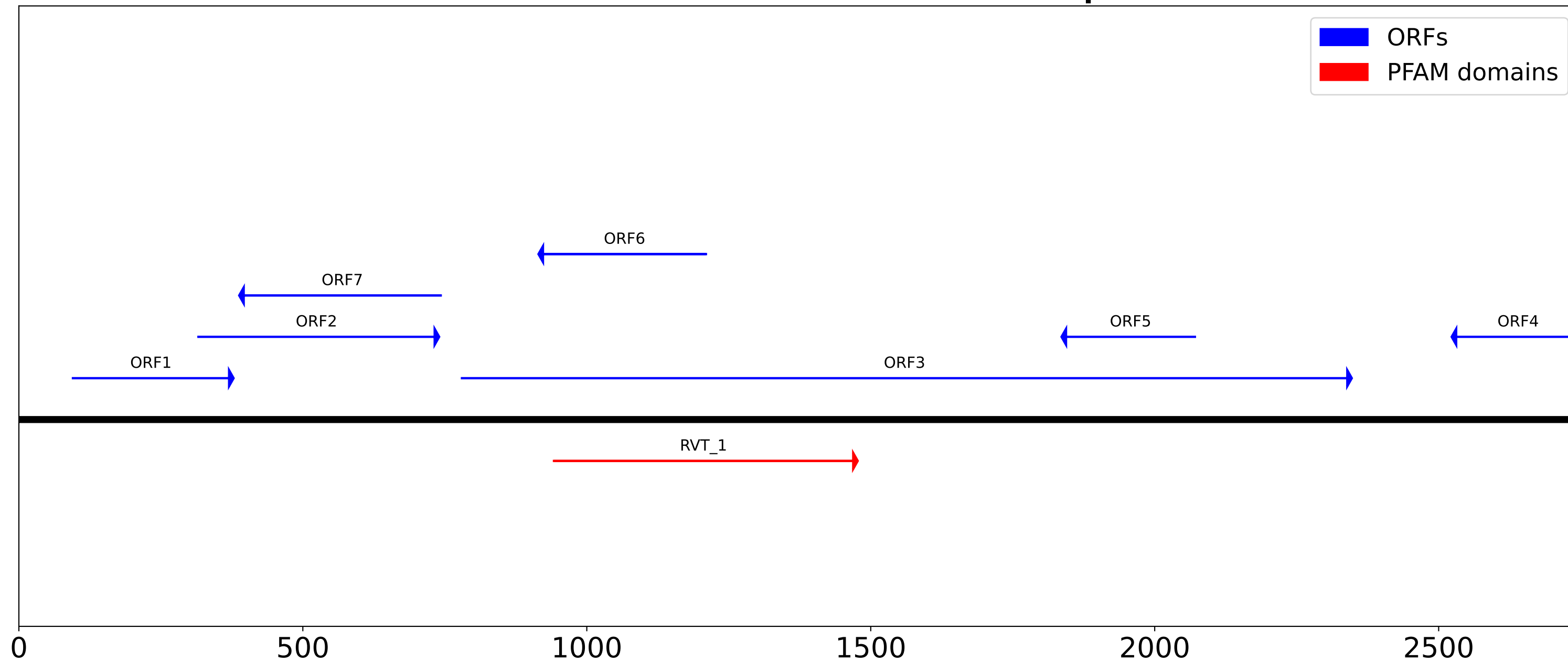


TE: rnd_1_family_832
size: 1024bp; fragments: 12539; full length: 415 (≥ 921.6 bp)

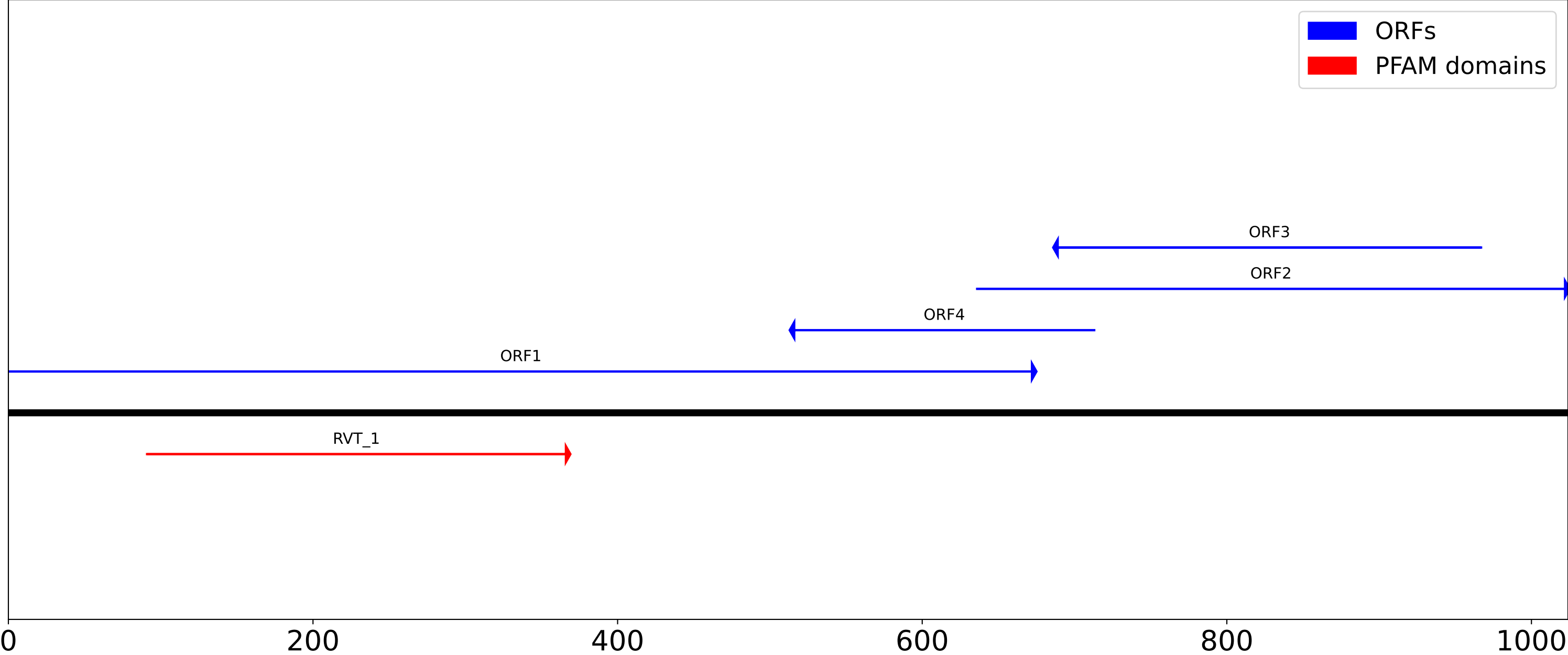
Before TEtrimmer 1024 bp



After TEtrimmer ORF and PFAM domain plot



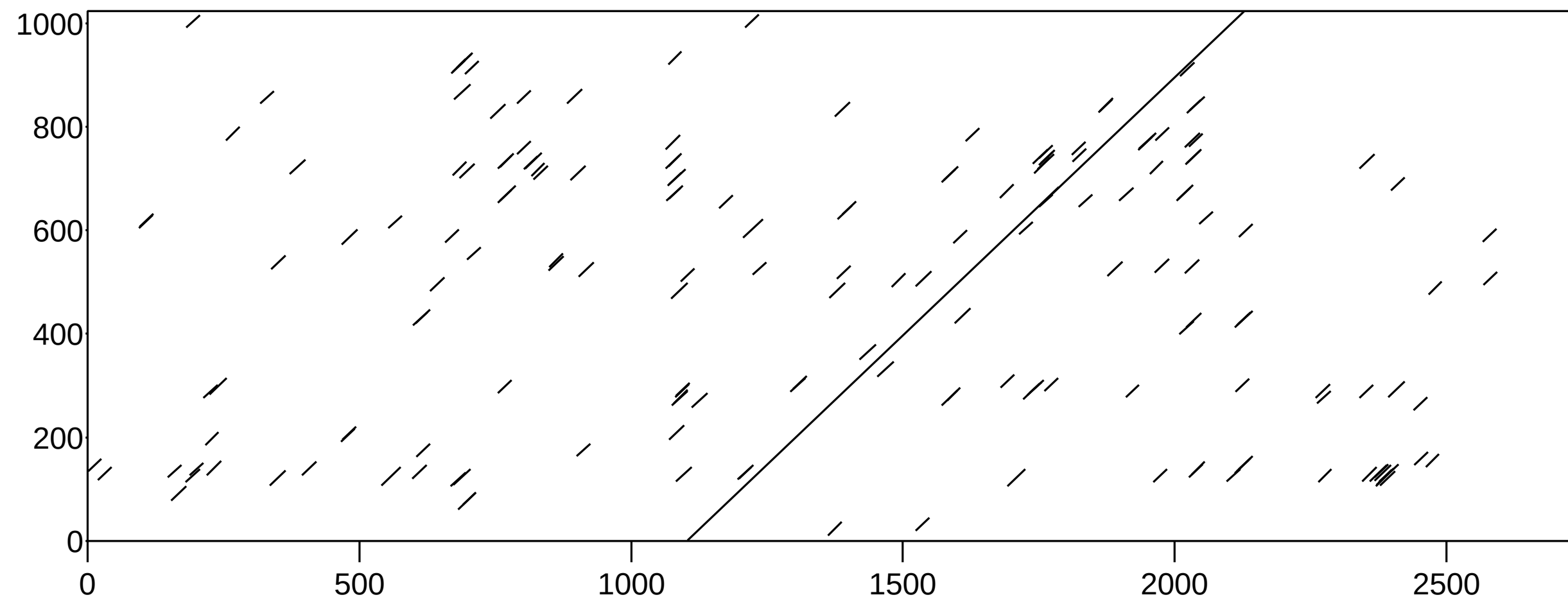
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)